

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Krona

RRID:SCR_012785

Type: Tool

Proper Citation

Krona (RRID:SCR_012785)

Resource Information

URL: <http://sourceforge.net/p/krona/home/krona/>

Proper Citation: Krona (RRID:SCR_012785)

Description: Software that allows hierarchical data to be explored with zoomable pie charts.

Abbreviations: Krona

Synonyms: Krona - Hierarchical data browser

Resource Type: software resource

Defining Citation: [PMID:21961884](#)

Keywords: bio.tools

Resource Name: Krona

Resource ID: SCR_012785

Alternate IDs: OMICS_01498, biotools:krona

Alternate URLs: <https://bio.tools/krona>

Record Creation Time: 20220129T080312+0000

Record Last Update: 20260905T132727+0000

Ratings and Alerts

No rating or validation information has been found for Krona.

No alerts have been found for Krona.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 97 mentions in open access literature.

Listed below are recent publications. The full list is available at [Kravitz](#) .

Zou X, et al. (2026) Introducing unprocessed oil-tea waste leads to imbalance of microbiome and disease spread for near-natural cultivation of shorthaired antelope. *Environmental microbiome*, 21(1).

Bie??o K, et al. (2026) Exploring the *Tenebrio molitor* gut microbiota response to LDPE and PET: putative genetic indicators and methodological insights. *Frontiers in microbiology*, 17, 1746922.

Sontigun N, et al. (2026) Epidemiology and Antimicrobial-Resistant Genes of Family Staphylococcaceae in *Musca domestica*: Case Studies from Chicken Farm, Pig Farms, and Residential Areas in Southern Thailand. *Insects*, 17(5).

Shulga EY, et al. (2026) Biotechnological Potential and Metabolic Diversity of Lignin-Degrading Bacteria from Decaying *Tilia cordata* Wood. *Microorganisms*, 14(2).

Xu H, et al. (2026) Pear scion genotypes shape rhizosphere fungal communities in association with photosynthetic sugar metabolism. *BMC plant biology*, 26(1).

Kim M, et al. (2026) Assessing the temporal dynamics of airborne insect communities through DNA metabarcoding of aerial trap catches. *Scientific reports*, 16(1).

Wen J, et al. (2026) Yeast enzyme hydrolysis slurry supplementation improves growth, intestinal health, and metabolic responses in juvenile largemouth bass (*Micropterus salmoides*) fed soybean meal-based diets with partial fishmeal replacement. *Journal of animal science and biotechnology*, 17(1).

Sun Q, et al. (2026) *Akkermansia muciniphila* drives viscero-visceral crosstalk via 5-HT_{3A}R-mediated sensitization of dichotomizing gut-bladder neurons. *Experimental & molecular medicine*, 58(5), 1608.

Diana L, et al. (2025) On-top arginine supplementation during lactation affects milk composition, performance, and intestinal bacterial and viral microbial community of sows and their piglets. *Journal of animal science*, 103.

Dimattia LM, et al. (2025) An ice inhabiting bdelloid rotifer from North America. *Extremophiles : life under extreme conditions*, 29(2), 30.

Trunfio M, et al. (2025) The cerebrospinal fluid virome in people with HIV: links to neuroinflammation and cognition. *bioRxiv : the preprint server for biology*.

Zhou M, et al. (2025) Microbial diagnostic features identified across populations possess potential antitumor properties in breast cancer. *mSystems*, 10(7), e0027125.

Zhang R, et al. (2025) Feeding probiotics-fermented distiller's grains diets increases rumen enzyme activities and glycerophospholipid levels in finishing cattle by modulating rumen microbiota. *Microbiome*, 13(1), 137.

Nasr E, et al. (2025) PathoGFAIR: a collection of FAIR and adaptable (meta)genomics workflows for (foodborne) pathogens detection and tracking. *GigaScience*, 14.

Gao Y, et al. (2025) Influence of highway construction of alpine grassland area in Gannan on soil properties and microbiota. *Scientific reports*, 15(1), 22964.

Morni MA, et al. (2025) Gut microbiome community profiling of Bornean bats with different feeding guilds. *Animal microbiome*, 7(1), 21.

Ma Z, et al. (2025) Apple replant disease: unraveling the fungal enigma hidden in the rhizosphere. *Stress biology*, 5(1), 71.

Han Y, et al. (2025) Influence of antibacterial peptides in drinking water on growth and intestinal health of European meat pigeon. *Poultry science*, 104(10), 105636.

Guo X, et al. (2025) *Klebsiella* predominance in peripancreatic microbial spectrum is associated with the severity of infected pancreatic necrosis. *BMC infectious diseases*, 25(1), 882.

Zhang R, et al. (2025) Multi-omics analyses reveal fecal microbial community and metabolic alterations in finishing cattle fed probiotics-fermented distiller's grains diets. *Microbiology spectrum*, 13(5), e0072124.